

- The female *Aedes aegypti* mosquito is the primary vector of the 2014 CHIKV outbreak in the Caribbean.
- Three genotypes of Chikungunya virus (CHIKV) exist which include the East Central South African, West African, and Asian Lineages.
- The Caribbean outbreak is associated with the Asian Lineage.
- The probability of the establishment of endemic Chikungunya in the Caribbean is high.
- Consequences of global travel and tourism within the Caribbean region will accelerate the continued introduction of CHIKV to the Americas and other countries.

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